

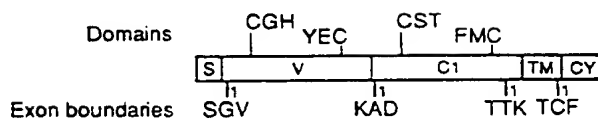
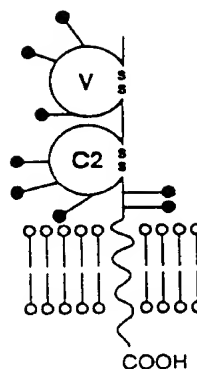
EX-4

Molecular weights
Polypeptides 30048

SDS PAGE
reduced 60 kD
unreduced 60 kD

Carbohydrate
N-linked sites 8
O-linked unknown

Human gene location and size
3q13.3-3q21; 32 kb¹



Tissue distribution

Present on a subset of B cells *in vivo* and the majority of B cells activated *in vitro*. Red blood cells, granulocytes, monocytes, resting or activated T cells, thymocytes and platelets do not express B7². The antigen is expressed by HTLV-I transformed T cells³.

Structure

The extracellular domain contains two IgSF domains which are highly glycosylated⁴. The sequence of the transmembrane domain is unusual in containing 3 cysteine residues that might be covalently modified or participate in intermolecular interactions⁴ although there is no evidence for this. The cytoplasmic domain has a preponderance (9/19) of arginine residues and contains a potential site for calmodulin-dependent phosphorylation (RRES)⁴.

Function

B7 is the ligand for the CD28⁵ and CTLA-4⁶ glycoproteins. Cells transfected with either human⁷ or murine⁸ B7 genes supply co-stimulatory signals to human T cells, suggesting that the CD28 binding site is conserved⁸. The antigen is not expressed on resting B cells but is strongly upregulated on B cells activated with a variety of agents, including the Epstein-Barr virus², cross-linking anti-IgM², anti-CD45 and anti-MHC Class II mAbs⁹, IL2 and IL4¹⁰. MAbs to B7 block the differentiation of B cells into antibody secreting cells¹¹ and the alloactivation of T cells⁹.

Comments

This antigen is not related to a mouse antigen called B7 and to avoid confusion the latter is being called B7(2).

Database accession numbers

	PIR	SWISSPROT	EMBL/GENBANK	REFERENCE
Human			M27533	4
Mouse			X60958	8

Amino acid sequence of human B7

MGHTRROGTS	PSKCPYLNFF	QLLVLA	-1
GLSHFCSGVI	HVTKEVKVA	TLSCGHNVS	50
MSGDMNIWPE	YKNRTIFDIT	NNLSIVILAL	100
KREHLAEVTL	SVKADFPTPS	ISDFEIPSTN	150
ENGEELNAIN	TTVSODPETE	LYAVSSKLD	200
QTFNWNTTKQ	EHFPDNLPS	WAITLISVNG	250
NERLRRESVR	PV		262

References

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